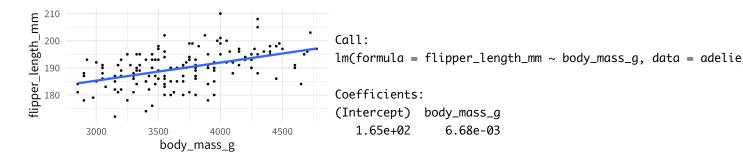
NOTES 24: INFERENCE FOR REGRESSION

Stat 120 | Fall 2025 Prof Amanda Luby

When we fit linear regression models earlier in this class, we used them to describe the relationship between the variables and interpreted the slope and the intercept as descriptions of the data. Now, we'd like to understand what the regression model can tell us about the relationship of the variables in the population.

We're going to return to the palmerpenguins dataset from earlier in the class and restrict our analysis today to Adelie penguins. We we want to predict flipper_length_mm using body_mass_g.



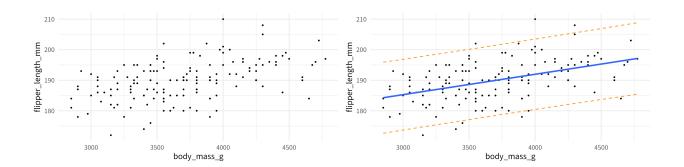
Warm up: Write out the regression equation and provide an interpretation for the slope



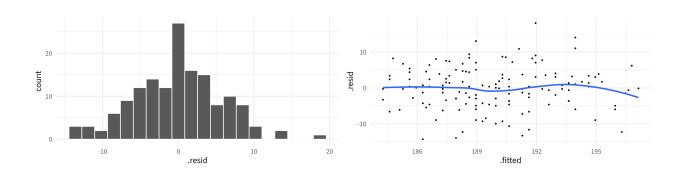
The **population model** for the regression line is:

Idea:

To check if this population model is reasonable, use the "LINE" mnemonic:



- 1. **L**_____
- 2. **I**_____
- 3. **N**_____
- 4. **E**_____



i Note

The **sampling distribution** for b_1 is:

summary(adelie_lm)

Call:

lm(formula = flipper_length_mm ~ body_mass_g, data = adelie)

Residuals:

Coefficients:

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 5.8 on 149 degrees of freedom

(1 observation deleted due to missingness)

Multiple R-squared: 0.219, Adjusted R-squared: 0.214 F-statistic: 41.8 on 1 and 149 DF, p-value: 1.34e-09

Note

CLT-based confidence interval for β_1 :

$$b_1 \pm t_{n-2}^* \times SE_{b_1}$$

Note

CLT-based test for $H_0: \beta_1 = 0$:

$$t_{n-2} = \frac{b_1 - 0}{SE_{b_1}}$$

Example: Using the summary() output, along with the sampling distribution information above, how would you compute a 95% confidence interval for β_1 ?

1 Inference for Predictions (Ch 9.3)

Example: We measure a new Adelie penguin that weighs 4000g. What is (a) the predicted *mean* flipper length among all 4000g penguins and (b) the predicted flipper length for *any* 4000g penguin?

```
augment(adelie_lm, newdata = tibble(body_mass_g = 4000))
# A tibble: 1 x 2
  body_mass_g .fitted
        <dbl>
                <dbl>
         4000
                 192.
1
augment(adelie_lm, newdata = tibble(body_mass_g = 4000), interval = "confidence")
# A tibble: 1 x 4
  body_mass_g .fitted .lower .upper
                <dbl> <dbl> <dbl>
1
         4000
                 192.
                        191.
                               193.
augment(adelie_lm, newdata = tibble(body_mass_g = 4000), interval = "prediction")
# A tibble: 1 x 4
  body_mass_g .fitted .lower .upper
               <dbl> <dbl> <dbl>
        <dbl>
         4000
                 192.
                               203.
1
                        180.
```

We usually rely on R to make confidence and prediction intervals, but the formulas for these intervals are:

Note ${\sf A} ____ {\sf interval for the mean response} \ {\sf when the predictor} \ {\sf is} \ {\sf x}^* \ {\sf is} :$

$$\hat{y} \pm t^*_{n-2} \times s_\epsilon \sqrt{\frac{1}{n} + \frac{(x^* - \bar{x})^2}{(n-1)s_x^2}}$$

Note $\hat{y}\pm t_{n-2}^*\times s_\epsilon\sqrt{1+\frac{1}{n}+\frac{(x^*-\bar{x})^2}{(n-1)s_x^2}}$